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Please amend the paragraph on page 31, lines 9 to 17 as follows:

Figure 14 is a graphical representation of a pairwise sequence alignment of the ERECTA proteins isolated from Arabidopsis (SEQ ID NO: 2), maize (SEQ ID NO: 45), rice (SEQ ID NO: 3), Sorghum (SEQ ID NO: 5) and wheat (SEQ ID NO: 20). The alignment was performed using CLUSTALW multiple sequence alignment tool. Residues that are conserved between all species are indicated by asterisks (*). Conservation of the groups STA (SEQ ID NO: 46) NEQK (SEQ ID NO: 47) NHQK (SEQ ID NO: 48) NDEQ (SEQ ID NO: 49) QHRK (SEQ ID NO: 50) MILV (SEQ ID NO: 51) MILF (SEQ ID NO: 52) HY (SEQ ID NO: 53) or FYW (SEQ ID NO: 54) is indicated by “:”. Conservation of the groups CSA (SEQ ID NO: 55) ATV (SEQ ID NO: 56) SAG (SEQ ID NO: 57) STNK (SEQ ID NO: 58) STPA (SEQ ID NO: 59) SGND (SEQ ID NO: 60) SNDEQK (SEQ ID NO: 61) NDEQHK (SEQ ID NO: 62) NEQHRK (SEQ ID NO: 63) FVLIM (SEQ ID NO: 64) HFY (SEQ ID NO: 65) is indicated by “.”. Gaps are indicated by dashes “-”.

After the Abstract of the Disclosure and before the Figures, please replace the paper copy of the Sequence Listing currently entered with the paper copy of the substitute Sequence Listing attached hereto as **Exhibit A**.